RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _

Source:

Date Processed by STIC:

ENTERED



IFW16

RAW SEQUENCE LISTING DATE: 04/27/2007
PATENT APPLICATION: US/10/734,661C TIME: 11:40:55

Input Set : A:\81408-4400 sequence listing.txt
Output Set: N:\CRF4\04272007\J734661C.raw

```
3 <110> APPLICANT: Yayon, Avner
              Rom, Eran
      5
              Thomassen-Wolf, Elisabeth
             Borges, Eric
      8 <120> TITLE OF INVENTION: ANTIBODIES THAT BLOCK RECEPTOR PROTEIN TYROSINE
KINASE ACTIVATION,
           . METHODS OF SCREENING AND USES THEREOF
     11 <130> FILE REFERENCE: 81408-4400
     13 <140> CURRENT APPLICATION NUMBER: US 10/734,661C
     14 <141> CURRENT FILING DATE: 2003-12-15
     16 <150> PRIOR APPLICATION NUMBER: US 60/299,187
     17 <151> PRIOR FILING DATE: 2001-06-20
     19 <150> PRIOR APPLICATION NUMBER: PCT/IL02/00494
     20 <151> PRIOR FILING DATE: 2002-06-20
     22 <160> NUMBER OF SEQ ID NOS: 106
     24 <170> SOFTWARE: PatentIn version 3.2
     26 <210> SEQ ID NO: 1
     27 <211> LENGTH: 806
     28 <212> TYPE: PRT
     29 <213> ORGANISM: Homo sapiens
     31 <300> PUBLICATION INFORMATION:
     32 <308> DATABASE ACCESSION NO: np 000133
     33 <309> DATABASE ENTRY DATE: 2001-02-21
     34 <313> RELEVANT RESIDUES: (1)..(806)
     36 <400> SEQUENCE: 1
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     46 Gly Arg Ala Ala Glu Val Pro Gly Pro Glu Pro Gly Gln Gln Glu Gln
     47
               35
                                    40
     50 Leu Val Phe Gly Ser Gly Asp Ala Val Glu Leu Ser Cys Pro Pro
                                55
     54 Gly Gly Pro Met Gly Pro Thr Val Trp Val Lys Asp Gly Thr Gly
                            70
                                                75
    58 Leu Val Pro Ser Glu Arg Val Leu Val Gly Pro Gln Arg Leu Gln Val
                        85
                                            90
    62 Leu Asn Ala Ser His Glu Asp Ser Gly Ala Tyr Ser Cys Arg Gln Arg
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     66 Leu Thr Gln Arg Val Leu Cys His Phe Ser Val Arg Val Thr Asp Ala
                                    120
     70 Pro Ser Ser Gly Asp Asp Glu Asp Glu Asp Glu Ala Glu Asp Thr
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74 Gly Val Asp Thr Gly Ala Pro Tyr Trp Thr Arg Pro Glu Arg Met Asp

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/734,661C

DATE: 04/27/2007 TIME: 11:40:55

75	145					150					155					160
		Lys	Leu	Leu	Ala		Pro	Ala	Ala	Asn	Thr	Val	Arq	Phe	Arq	
79	•	•			165					170					175	1
82	Pro	Ala	Ala	Gly	Asn	Pro	Thr	Pro	Ser	Ile	Ser	Trp	Leu	Lys	Asn	Gly
83				180					185			_		190	•	
86 .	Arg	Glu	Phe	Arg	Gly	Glu	His	Arg	Ile	Gly	Gly	Ile	Lys	Leu	Arg	His
87			195					200					205			
90	Gln	Gln	Trp	Ser	Leu	Val	Met	Glu	Ser	Val	Val	Pro	Ser	Asp	Arg	Gly
91		210					215					220				
94 .	Asn	Tyr	Thr	Cys	Val	Val	Glu	Asn	Lys	Phe	Gly	Ser	Ile	Arg	Gln	Thr
95						230					235					240
	Tyr	Thr	Leu	Asp	Val	Leu	Glu	Arg	Ser	Pro	His	Arg	Pro	Ile	Leu	Gln
99					245					250					255	
	Ala	Gl	Let			Asr.	ı Glr	Thi			. Let	Gly	Ser	-		Glu
103				260					265				_	270		
		His		_	val	. Tyr	Ser			Glr	Pro	His			Trp	Leu
107			275			_		280				_	285			_
				. GI	ı Val	. Asr	_		Lys	val	. Gly		_	GT?	Thr	Pro
111	•	290			T		295			- 77-		300			· • • • • •	
	_		ını	val	. Let	. Lys 310		. Als	r GTZ	Ala			Thi	Asp	р гуг	Glu
	305		. 1751	Τ 01				. 7~~	. 17-1	The	315		7.4	. 77-	~1.	320 Glu
119		GIU	ı vaı	. пес	325		i nis	ASI	ı val	330		GIU	ASL	Alc	335	
		Thr	- Cvc	. T.A1			, Den	Ser	· T]_			Ser	· Wie	. Wie		Ala
123	- 7 -		. • , .	340		. Oly	7101.		345	_	1110	. DCI	1111	350		ALG
	Tre	Leu	ı Val			Pro	Ala	Glu			Leu	Val	Glu			Glu
127			355					360					365			
		Gly	Ser	Val	Tyr	Ala	Gly			Ser	Tyr	Glv			, Phe	Phe
131		370			•		375				•	380		. 4		
134	Leu	Phe	: Ile	Leu	ı Val	. Val	. Ala	Ala	\Val	Thr	Leu	Cys	Arg	Leu	. Arg	Ser
135	385					390)				395					400
	Pro	Pro	Lys	Lys	Gly	Leu	ı Gly	Ser	Pro	Thr	Val	His	Lys	: Ile	Ser	Arg
139					405					410					415	
	Phe	Pro	Leu	_		Gln	val	Ser			Ser	Asn	Ala			Ser
143	_	_		420			_		425				_	430		
	Ser	Asr			Leu	val	. Arg			Arg	Leu	Ser			Glu	Gly
147	D	mb -	435		3	77-7	0	440		. al	.	5	445			
151	PIO			Ala	ASI	val			Let	GIU	Leu			Asp	Pro	Lys
	T~~	450		60*	. 7.~~	. או	455		The		~1·	460			~1.	. ~1
	465		пеп	. ser	Arg	470		пес	1111	пеп	475		PLC	, пеп	GIY	Glu 480
			Dhe	ഭിച	. Gln			Mot	- Δ1 ≃	GI 11			Gla	, Tla	λαν	Lys
159	CIY	Cys		Оту	485		· val	1-1-6	. AIC	490		110	СТУ	116	495	_
	Asp	Aro	Ala	Ala			Val	Thr	· Val			Lvs	Met	Len		Asp
163		3		500					505			_,,		510	_	
	Asp	Ala	Thr			Asp	Leu	Ser			Val	Ser	Glu			Met
167	-		515					520					525			
	Met	Lys	Met	Ile	Gly	Lys	His	Lys	Asn	Ile	Ile	Asn	Leu	Leu	Gly	Ala
171		530			_	_	535					540			-	

RAW SEQUENCE LISTING DATE: 04/27/2007 PATENT APPLICATION: US/10/734,661C TIME: 11:40:55

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174 Cys Thr Gln Gly Gly Pro Leu Tyr Val Leu Val Glu Tyr Ala Ala Lys
175 545
178 Gly Asn Leu Arg Glu Phe Leu Arg Ala Arg Arg Pro Pro Gly Leu Asp
179
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182 Tyr Ser Phe Asp Thr Cys Lys Pro Pro Glu Glu Gln Leu Thr Phe Lys
               580
                                    585
186 Asp Leu Val Ser Cys Ala Tyr Gln Val Ala Arg Gly Met Glu Tyr Leu
     595
                                600
190 Ala Ser Gln Lys Cys Ile His Arg Asp Leu Ala Ala Arg Asn Val Leu
      610
                            615
                                                620
194 Val Thr Glu Asp Asn Val Met Lys Ile Ala Asp Phe Gly Leu Ala Arg
                        630
                                            635
198 Asp Val His Asn Leu Asp Tyr Tyr Lys Lys Thr Thr Asn Gly Arg Leu
                    645
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202 Pro Val Lys Trp Met Ala Pro Glu Ala Leu Phe Asp Arg Val Tyr Thr
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                                    665
206 His Gln Ser Asp Val Trp Ser Phe Gly Val Leu Leu Trp Glu Ile Phe
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210 Thr Leu Gly Gly Ser Pro Tyr Pro Gly Ile Pro Val Glu Glu Leu Phe
                            695
214 Lys Leu Leu Lys Glu Gly His Arg Met Asp Lys Pro Ala Asn Cys Thr
215 705
                        710
                                            715
218 His Asp Leu Tyr Met Ile Met Arg Glu Cys Trp His Ala Ala Pro Ser
                                        730
222 Gln Arg Pro Thr Phe Lys Gln Leu Val Glu Asp Leu Asp Arg Val Leu
                740
                                    745
226 Thr Val Thr Ser Thr Asp Glu Tyr Leu Asp Leu Ser Ala Pro Phe Glu
            755
                                760
230 Gln Tyr Ser Pro Gly Gly Gln Asp Thr Pro Ser Ser Ser Ser Gly
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234 Asp Asp Ser Val Phe Ala His Asp Leu Leu Pro Pro Ala Pro Pro Ser
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238 Ser Gly Gly Ser Arg Thr
242 <210> SEQ ID NO: 2
243 <211> LENGTH: 32
244 <212> TYPE: DNA
245 <213> ORGANISM: Artificial Sequence
247 <220> FEATURE:
248 <223> OTHER INFORMATION: artificial primer
250 <400> SEQUENCE: 2
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255 <211> LENGTH: 55
256 <212> TYPE: DNA
257 <213> ORGANISM: Artificial Sequence
259 <220> FEATURE:
260 <223> OTHER INFORMATION: artificial primer
262 <400> SEQUENCE: 3
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RAW SEQUENCE LISTING DATE: 04/27/2007
PATENT APPLICATION: US/10/734,661C TIME: 11:40:55

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266 <210> SEQ ID NO: 4
267 <211> LENGTH: 1147
268 <212> TYPE: DNA
269 <213> ORGANISM: Homo sapiens
271 <300> PUBLICATION INFORMATION:
272 <308> DATABASE ACCESSION NO: m58051
273 <309> DATABASE ENTRY DATE: 1994-11-08
274 <313> RELEVANT RESIDUES: (1)..(1147)
276 <400> SEQUENCE: 4
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279 tegegetetg egtggeegtg gecategtgg eeggegeete eteggagtee ttggggaegg
                                                                        120
281 agcagcgcgt cgtggggcga gcggcagaag tcccgggccc agagcccggc cagcaggagc
                                                                        180
240
285 ccatggggcc cactgtctgg gtcaaggatg gcacagggct ggtgccctcg gagcgtgtcc
                                                                        300
287 tggtggggcc ccagcggctg caggtgctga atgcctccca cgaggactcc ggggcctaca
                                                                        360
289 gctgccggca gcggctcacg cagcgcgtac tgtgccactt cagtgtgcgg gtgacagacg
                                                                        420
291 ctccatcctc gggagatgac gaagacgggg aggacgaggc tgaggacaca ggtgtggaca
                                                                        480
293 caggggcccc ttactggaca cggcccgagc ggatggacaa gaagctgctg gccgtgccgg
                                                                        540
295 ccgccaacac cgtccgcttc cgctgcccag ccgctggcaa ccccactccc tccatctcct
                                                                        600
297 ggctgaagaa cggcagggag ttccgcggcg agcaccgcat tggaggcatc aagctgcggc
                                                                        660
299 atcagcagtg gagcctggtc atggaaagcg tggtgccctc ggaccgcggc aactacacct
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301 gegtegtgga gaacaagttt ggeageatee ggeagaegta caegetggae gtgetggage
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303 gctccccgca ccggcccatc ctgcaggcgg ggctgccggc caaccagacg gcggtgctgg
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305 gcagcgacgt ggagttccac tgcaaggtgt acagtgacgc acagccccac atccagtggc
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307 tcaagcacgt ggaggtgaac ggcagcaagg tgggcccgga cggcacaccc tacgttaccg
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309 tgctcaagac ggcgggcgct aacaccaccg acaaggagct agaggttctc tccttgcaca
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311 acgtcacctt tgaggacgcc ggggagtaca cctgcctggc gggcaattct attgggtttt
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315 aggcggg
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319 <211> LENGTH: 5695
320 <212> TYPE: DNA/
321 <213> ORGANISM (EXPRESSION VECTOR pCEP-PU/AC7
323 <400> SEQUENCE: 5
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328 cgagcaaaat ttaagctaca acaaggcaag gcttgaccga caattgcatg aagaatctgc
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330 ttagggttag gegttttgeg etgettegeg atgtaeggge cagatataeg egttgaeatt
                                                                        240
332 gattattgac tagttattaa tagtaatcaa ttacggggtc attagttcat agcccatata
334 tggagttccg cgttacataa cttacggtaa atggcccgcc tggctgaccg cccaacgacc
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336 cccgcccatt gacgtcaata atgacgtatg ttcccatagt aacgccaata gggactttcc
338 attgacgtca atgggtggac tatttacggt aaactgccca cttggcagta catcaagtqt
340 atcatatgcc aagtacgccc cctattgacg tcaatgacgg taaatggccc gcctggcatt
342 atgcccagta catgacctta tgggactttc ctacttggca gtacatctac gtattagtca
                                                                        600
344 tegetattae catggtgatg eggttttgge agtacateaa tgggegtgga tageggtttg
                                                                        660
346 actcacgggg atttccaagt ctccacccca ttgacgtcaa tgggagtttg ttttggcacc
                                                                        720
348 aaaatcaacg ggactttcca aaatgtcgta acaactccgc cccattgacg caaatgggcg
                                                                        780
350 gtaggegtgt aeggtgggag gtetatataa geagagetet etggetaaet agagaaeeea
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RAW SEQUENCE LISTING DATE: 04/27/2007 PATENT APPLICATION: US/10/734,661C TIME: 11:40:55

352	ctgcttactg	gcttatcgaa	attaatacqa	ctcactatag	ggagacccaa	gctggctagc	900
	gtttaaactt						960
	gagatcccga						1020
	ctgaactcct						1080
	tgatctcccg						1140
	aggtcaagtt						1200
	gggaggagca						1260
	actggctgaa						1320
	tcgagaaaac						1380
	ccccatcccg					_	1440
	tctatcccag						1500
	agaccacgcc						1560
	tggacaagag						1620
	tgcacaacca						1680
	ccgtttaaac						1740
	gccctcccc						1800
	aaaatgagga						1860
	tggggcagga						1920
	tgggctctat						1980
	cgccctgtag						2040
	cacttgccag						2100
	tcgccggctt						2160
	ctttacggca						2220
	cgccctgata						2280
	tcttgttcca						2340
	ggattttggg						2400
	cgaattaatt						2460
	caggcagaag						2520
	caggeteece						2580
	tcccgcccct.						2640
	cccatggctg						2700
	tattccagaa						2760
	gagcttgtat						2820
	gtatatcggc						2880
	tgccgttccg						2940
	gctcgggttc						3000
	gaccctgttc						3060
	gtgggtgcgc						3120
	ccgggacgcc						3180
	cgccctgcgc						3240
	cgtgctacga						3300
	tttccgggac						3360
	ccaccccaac						3420
	tttcacaaat						3480
	tgtatcttat						3540
	atagctgttt						3600
	aagcataaag						3660
	gcgctcactg						3720
				tgggcgctct			3780

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/734,661C

DATE: 04/27/2007 TIME: 11:40:56

Input Set : A:\81408-4400 sequence listing.txt
Output Set: N:\CRF4\04272007\J734661C.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:54; N Pos. 253,254,255

Seq#:56; N Pos. 256,257,258

Seq#:70; N Pos. 1,2,3

Seq#:74; N Pos. 1,2,3 Seq#:81; N Pos. 1,2,3

Seq#:83; N Pos. 1,2,3

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/734,661C

DATE: 04/27/2007 TIME: 11:40:56

Input Set : A:\81408-4400 sequence listing.txt
Output Set: N:\CRF4\04272007\J734661C.raw

L:1612 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54 after pos.:240
L:1662 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56 after pos.:240
L:1968 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 after pos.:0
L:2064 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74 after pos.:0
L:2234 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:81 after pos.:0
L:2286 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:0